

OIPE 0570
083

CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date: 10/10/2001
Edited by: AN
Verified by: AN (STIC stat)

Serial Number: 09/924,340

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was wrapped down to the next line.

Edited a formal error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included: _____

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/username at end of file;
 page numbers throughout text; other invalid text, such as _____

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

Examiner: The above corrections must be communicated to the applicant in the first office
Action: DO NOT send a copy of this form.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/924,340

DATE: 10/10/2001
TIME: 18:15:39

Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

2 <110> APPLICANT: Bejanin, Stephane
3 Tanaka, Hiroaki
5 <120> TITLE OF INVENTION: HUMAN CDNAS AND PROTEINS AND USES THEREOF
7 <130> FILE REFERENCE: 91.US2.REG
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/924,340
10 <141> CURRENT FILING DATE: 2001-08-06
12 <150> PRIOR APPLICATION NUMBER: US 60/305,456
13 <151> PRIOR FILING DATE: 2001-07-13
15 <150> PRIOR APPLICATION NUMBER: US 60/302,277
16 <151> PRIOR FILING DATE: 2001-06-29
18 <150> PRIOR APPLICATION NUMBER: US 60/298,698
19 <151> PRIOR FILING DATE: 2001-06-15
21 <150> PRIOR APPLICATION NUMBER: US 60/293,574
22 <151> PRIOR FILING DATE: 2001-05-25
24 <160> NUMBER OF SEQ ID NOS: 112
26 <170> SOFTWARE: JPatent
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 2016
30 <212> TYPE: DNA
31 <213> ORGANISM: Homo sapiens
33 <220> FEATURE:
34 <221> NAME/KEY: 5'UTR
35 <222> LOCATION: 1..1434
37 <220> FEATURE:
38 <221> NAME/KEY: CDS
39 <222> LOCATION: 1435..1836
41 <220> FEATURE:
42 <221> NAME/KEY: 3'UTR
43 <222> LOCATION: 1837..2016
45 <220> FEATURE:
46 <221> NAME/KEY: polyA_signal
47 <222> LOCATION: 1965..1970
49 <220> FEATURE:
50 <221> NAME/KEY: polyA_site
51 <222> LOCATION: 2001..2016
53 <400> SEQUENCE: 1
54 aagggtcttc tgcatacata caccaaggaa aagccacatg aggacataac caggaagaga 60
55 gccatcacca agaaccggaa catgcggaca ccctgatctc ggacttctag ctttcagaac 120
56 cgttgcacca gttttgatga tcatctctt cccaaaccaag atggtgaaa aagaaaaac 180
57 gtggtaatc ttggagcaat ccgacaaggc atgaaacgct tccaatttct gttaaactgc 240
58 tgtgagccag ggacaattcc tggatgcattt atccttagcag ctgccttggaa tctactatgc 300
59 ggcattcttc tgattcattt ttctccattt gtgtgtttt tctctgtat gtgaatccat 360
60 ccctatccat tatgtcatgc ctccatctt tgctgcttct tcagattgca ctgagccata 420
61 agaggaagcc cctgtgggtgg ccagagcagc cttgttcctg gaatgtgctc gttttgttca 480
62 cgcgtcaac cgtggcaact ggccagatgt gatgaaaggg caccacgtga acatcaccaa 540
63 gaaaggactt tccggggac gtcctccat tggatgcac aagcggaaacc agaagctgca 600
64 gtggaatgca gccaagctct tctaccaatg gggagacaag gaaaaaaaggtaaaa 660

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Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

65 agaaattca agaggacaa gtttctgcta attttagaca gagctgaaca taaacacaca 720
 66 taaagagggt ccatatattc ctctttctt aaagattact tggataact gttacaattt 780
 67 ccgtaataa ttcagctgaa tggcttacc aatgtgccta ccaactaagg caattggcgt 840
 68 ccgattgaat gagctgtcc acggggaaag tgagagccca gccaacctgc tgggtctcat 900
 69 ttacgatgaa gagaccaaga ggagacttag aaaggaggat gaggaggaag actttttaga 960
 70 tgacattcca cttcaagtc aatacacagc tcattgcata tttaaagct gattatgtg 1020
 71 caagcaactt tcgggctgga aattctacag aagcttgcata tttccattct tgatgagagg 1080
 72 caaagtcccc ggcaacaaaat taactcagga gagaaaatgg tttcctgaa aaaaacgata 1140
 73 gcttaataat ctacagaaaag accgttaattt ccacctattt tcaatgaaa tcgtgaaaaa 1200
 74 cacatttggc ctagagctga aacaacttca ctgcctcaa aacagcaaga cagacatccc 1260
 75 tcataaaatg aactgacaga atttttagt ctccaaatct agtcaactgc catacacata 1320
 76 gtctaaatct gattgaatag cagctgtagaa atcttgcgaa attacttccc atttctgtt 1380
 77 tcgttaaaag gtactgtgaa cccctctaaa tgcgggtgccc ccttgcctt gaag atg 1437
 78 Met
 79 1
 80 gca gca tgt cag ctt ctt ctg gag att acc acc ttc ctg cga gag acc 1485
 81 Ala Ala Cys Gln Leu Leu Leu Glu Ile Thr Thr Phe Leu Arg Glu Thr
 82 5 10 15
 83 ttt tct tgc ctg ccc aga cct cgc act gag cct ctg gtg gct tca acg 1533
 84 Phe Ser Cys Leu Pro Arg Pro Arg Thr Glu Pro Léu Val Ala Ser Thr
 85 20 25 30
 86 gac cac acc aaa atg cca tct caa atg gaa cac gcc atg gaa acc atg 1581
 87 Asp His Thr Lys Met Pro Ser Gln Met Glu His Ala Met Glu Thr Met
 88 35 40 45
 89 atg ttt aca ttt cac aaa ttc gct ggg gat aaa ggc tac tta aca aag 1629
 90 Met Phe Thr Phe His Lys Phe Ala Gly Asp Lys Gly Tyr Leu Thr Lys
 91 50 55 60 65
 92 gag gac ctg aga gta ctc atg gaa aag gag ttc cct gga ttt ttg gaa 1677
 93 Glu Asp Leu Arg Val Leu Met Glu Lys Glu Phe Pro Gly Phe Leu Glu
 94 70 75 80
 95 aat caa aaa gac cct ctg gct gtg gac aaa ata atg aag gac ctg gac 1725
 96 Asn Gln Lys Asp Pro Leu Ala Val Asp Lys Ile Met Lys Asp Leu Asp
 97 85 90 95
 98 cag tgt aga gat ggc aaa gtg ggc ttc cag agc ttc ttt tcc cta att 1773
 99 Gln Cys Arg Asp Gly Lys Val Gly Phe Gin Ser Phe Phe Ser Leu Ile
 100 100 105 110
 101 ggc ggc ctc acc att gca tgc aat gac tat ttt gta gta cac atg aag 1821
 102 Ala Gly Leu Thr Ile Ala Cys Asn Asp Tyr Phe Val Val His Met Lys
 103 115 120 125
 104 cag aag gga aag aag taggcagaaa tgagcagttc gtcctccct gataagagtt 1876
 105 Gln Lys Gly Lys Lys
 106 130
 107 gtcggcttaag gaatctgccc cacagttcc cccatagaag gatttcatga 1936
 108 gcagatcagg acacttagca aatgtaaaaa taaaatctaa ctctcattt acaagcagag 1996
 109 aaagaaaaaa aaaaaaaaaat
 111 <210> SEQ ID NO: 2
 112 <211> LENGTH: 134
 113 <212> TYPE: PRT
 114 <213> ORGANISM: Homo sapiens

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Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

116 <400> SEQUENCE: 2
117 Met Ala Ala Cys Gln Leu Leu Leu Glu Ile Thr Thr Phe Leu Arg Glu
118 1 5 10 15
119 Thr Phe Ser Cys Leu Pro Arg Pro Arg Thr Glu Pro Leu Val Ala Ser
120 20 25 30
121 Thr Asp His Thr Lys Met Pro Ser Gln Met Glu His Ala Met Glu Thr
122 35 40 45
123 Met Met Phe Thr Phe His Lys Phe Ala Gly Asp Lys Gly Tyr Leu Thr
124 50 55 60
125 Lys Glu Asp Leu Arg Val Leu Met Glu Lys Glu Phe Pro Gly Phe Leu
126 65 70 75 80
127 Glu Asn Gln Lys Asp Pro Leu Ala Val Asp Lys Ile Met Lys Asp Leu
128 85 90 95
129 Asp Gln Cys Arg Asp Gly Lys Val Gly Phe Gln Ser Phe Phe Ser Leu
130 100 105 110
131 Ile Ala Gly Leu Thr Ile Ala Cys Asn Asp Tyr Phe Val Val His Met
132 115 120 125
133 Lys Gln Lys Gly Lys Lys
134 130
136 <210> SEQ ID NO: 3
137 <211> LENGTH: 1081
138 <212> TYPE: DNA
139 <213> ORGANISM: Homo sapiens
141 <220> FEATURE:
142 <221> NAME/KEY: 5'UTR
143 <222> LOCATION: 1..38
145 <220> FEATURE:
146 <221> NAME/KEY: CDS
147 <222> LOCATION: 39..917
149 <220> FEATURE:
150 <221> NAME/KEY: 3'UTR
151 <222> LOCATION: 918..1081
153 <220> FEATURE:
154 <221> NAME/KEY: polyA_signal
155 <222> LOCATION: 1045..1050
157 <220> FEATURE:
158 <221> NAME/KEY: polyA_site
159 <222> LOCATION: 1066..1081
161 <400> SEQUENCE: 3
162 gtccagcctg ttgctgtatgc tgccgtgcgg tacttgtc atg gag ctg gca ctg cgg 56
163 Met Glu Leu Ala Leu Arg -20
164 -25 -20 104
165 cgc tct ccc gtc ccg cgg tgg ttg ctg ctg ccg ctg ctg ggc
166 Arg Ser Pro Val Pro Arg Trp Leu Leu Leu Pro Leu Leu Gly -5
167 -15 -10 152
168 ctg aac gca gga gct gtc att gac tgg ccc aca gag gag ggc aag gaa
169 Leu Asn Ala Gly Ala Val Ile Asp Trp Pro Thr Glu Glu Gly Lys Glu 10
170 1 5 10 200
171 gta tgg gat tat gtg acg gtc cgc aag gat gcc tac atg ttc tgg tgg

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Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

172 Val Trp Asp Tyr Val Thr Val Arg Lys Asp Ala Tyr Met Phe Trp Trp		
173 15 20 25		248
174 ctc tat tat gcc acc aac tcc tgc aag aac ttc tca gaa ctt ccc ctg		
175 Leu Tyr Tyr Ala Thr Asn Ser Cys Lys Asn Phe Ser Glu Leu Pro Leu		
176 30 35 40 45		
177 gtc atg tgg ctt cag ggc ggt cca ggc ggt tct agc act gga ttt gga		296
178 Val Met Trp Leu Gln Gly Gly Pro Gly Gly Ser Ser Thr Gly Phe Gly		
179 50 55 60		
180 aac ttt gag gaa att ggg ccc ctt gac agt gat ctc aaa cca cgg aaa		344
181 Asn Phe Glu Glu Ile Gly Pro Leu Asp Ser Asp Leu Lys Pro Arg Lys		
182 65 70 75		
183 acc acc tgg ctc cag gct gcc agt ctc cta ttt gtg gat aat ccc gtg		392
184 Thr Thr Trp Leu Gln Ala Ala Ser Leu Leu Phe Val Asp Asn Pro Val		
185 80 85 90		
186 ggc act ggg ttc agt tat gtg aat ggt agt ggt gcc tat gcc aag gac		440
187 Gly Thr Gly Phe Ser Tyr Val Asn Gly Ser Gly Ala Tyr Ala Lys Asp		
188 95 100 105		
189 ctg gct atg gtg gct tca gac atg atg gtt ctc ctg aag acc ttc ttc		488
190 Leu Ala Met Val Ala Ser Asp Met Met Val Leu Leu Lys Thr Phe Phe		
191 110 115 120 125		
192 agt tgc cac aaa gaa ttc cag aca gtt cca ttc tac att ttc tca gag		536
193 Ser Cys His Lys Glu Phe Gln Thr Val Pro Phe Tyr Ile Phe Ser Glu		
194 130 135 140		
195 tcc tat gga gga aaa atg gca gct ggc att ggt cta gag ctt tat aag		584
196 Ser Tyr Gly Gly Lys Met Ala Ala Gly Ile Gly Leu Glu Leu Tyr Lys		
197 145 150 155		
198 gcc att cag cga ggg acc atc aag tgc aac ttt gcg ggg gtt gcc ttg		632
199 Ala Ile Gln Arg Gly Thr Ile Lys Cys Asn Phe Ala Gly Val Ala Leu		
200 160 165 170		
201 ggt gat tcc tgg atc tcc cct gtt gat tcg gtg ctc tcc tgg gga cct		
202 Gly Asp Ser Trp Ile Ser Pro Val Asp Ser Val Leu Ser Trp Gly Pro		
203 175 180 185		
204 tac ctg tac agc atg tct ctt ctc gaa gac aaa ggt ctg gca gag gtg		728
205 Tyr Leu Tyr Ser Met Ser Leu Leu Glu Asp Lys Gly Leu Ala Glu Val		
206 190 195 200 205		
207 tct aag gtt gca gag caa gta ctg aat gcc gta aat aag ggg ctc tac		776
208 Ser Lys Val Ala Glu Gln Val Leu Asn Ala Val Asn Lys Gly Leu Tyr		
209 210 215 220		
210 aga gag gcc aca gag ctg tgg ggg aaa gca gaa atg atc att gaa cag		824
211 Arg Glu Ala Thr Glu Leu Trp Gly Lys Ala Glu Met Ile Ile Glu Gln		
212 225 230 235		
213 gta aaa agg gga aac act cag agg cta gcc tgc ttg gct ttt tct ggt		872
214 Val Lys Arg Gly Asn Thr Gln Arg Leu Ala Cys Leu Ala Phe Ser Gly		
215 240 245 250		
216 ggg tac agg gcc cat ggt tgg tgt tgt caa act tgg agt cta cac		917
217 Gly Tyr Arg Ala His Gly Trp Cys Cys Gln Thr Trp Ser Leu His		
218 255 260 265		
219 tgaggctccc cacatatctg caaatgattt catgctggat aataaatctc ttgggtctaa		977
220 gcagtatgtt agtgatgtt tacagatgtca gaaaggccacc caggcctgca agacttgctt		1037

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Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

1081

221 gtccttcact aaatgtatgg attctattaa aaaaaaaaaaaa aaaa
 223 <210> SEQ ID NO: 4
 224 <211> LENGTH: 293
 225 <212> TYPE: PRT
 226 <213> ORGANISM: Homo sapiens
 228 <220> FEATURE:
 229 <221> NAME/KEY: SIGNAL
 230 <222> LOCATION: 1..26
 232 <400> SEQUENCE: 4
 233 Met Glu Leu Ala Leu Arg Arg Ser Pro Val Pro Arg Trp Leu Leu Leu
 234 -25 -20 -15
 235 Leu Pro Leu Leu Leu Gly Leu Asn Ala Gly Ala Val Ile Asp Trp Pro
 236 -10 -5 1 5
 237 Thr Glu Glu Gly Lys Glu Val Trp Asp Tyr Val Thr Val Arg Lys Asp
 238 10 15 20
 239 Ala Tyr Met Phe Trp Trp Leu Tyr Tyr Ala Thr Asn Ser Cys Lys Asn
 240 25 30 35
 241 Phe Ser Glu Leu Pro Leu Val Met Trp Leu Gln Gly Pro Gly Gly
 242 40 45 50
 243 Ser Ser Thr Gly Phe Gly Asn Phe Glu Glu Ile Gly Pro Leu Asp Ser
 244 55 60 65 70
 245 Asp Leu Lys Pro Arg Lys Thr Thr Trp Leu Gln Ala Ala Ser Leu Leu
 246 75 80 85
 247 Phe Val Asp Asn Pro Val Gly Thr Gly Phe Ser Tyr Val Asn Gly Ser
 248 90 95 100
 249 Gly Ala Tyr Ala Lys Asp Leu Ala Met Val Ala Ser Asp Met Met Val
 250 105 110 115
 251 Leu Leu Lys Thr Phe Phe Ser Cys His Lys Glu Phe Gln Thr Val Pro
 252 120 125 130
 253 Phe Tyr Ile Phe Ser Glu Ser Tyr Gly Gly Lys Met Ala Ala Gly Ile
 254 135 140 145 150
 255 Gly Leu Glu Leu Tyr Lys Ala Ile Gln Arg Gly Thr Ile Lys Cys Asn
 256 155 160 165
 257 Phe Ala Gly Val Ala Leu Gly Asp Ser Trp Ile Ser Pro Val Asp Ser
 258 170 175 180
 259 Val Leu Ser Trp Gly Pro Tyr Leu Tyr Ser Met Ser Leu Leu Glu Asp
 260 185 190 195
 261 Lys Gly Leu Ala Glu Val Ser Lys Val Ala Glu Gln Val Leu Asn Ala
 262 200 205 210
 263 Val Asn Lys Gly Leu Tyr Arg Glu Ala Thr Glu Leu Trp Gly Lys Ala
 264 215 220 225 230
 265 Glu Met Ile Ile Glu Gln Val Lys Arg Gly Asn Thr Gln Arg Leu Ala
 266 235 240 245
 267 Cys Leu Ala Phe Ser Gly Gly Tyr Arg Ala His Gly Trp Cys Cys Gln
 268 250 255 260
 269 Thr Trp Ser Leu His
 270 265
 272 <210> SEQ ID NO: 5
 273 <211> LENGTH: 438

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

10/10/01

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/924,340

DATE: 10/10/2001
TIME: 18:15:40

Input Set : N:\jumbos\924340.txt
Output Set: N:\CRF3\10102001\I924340.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:1470 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27
L:1470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1491 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:27
L:1491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1537 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1551 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1592 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:29
L:1592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1655 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:4089 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:71
L:4089 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:4156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72